

Amendments to the Specification:

Please replace the original paper copy of the Sequence Listing with the substitute paper copy of the Sequence Listing filed herewith.

Please amend paragraph beginning at page 8, line 2, as follows:

Figure 1 (A) is a schematic illustration of the structure of wild type (replication-competent) MoMLV retrovirus; LTR= long terminal repeat. (B) is a schematic representation of glZD-GFP and glZD-hygro, showing the sizes of the IRES-transgene cassettes and the site of their insertion into the wild-type MLV genome. Arrows indicate location of *NheI* sites used to digest DNA for Southern hybridization analysis, and wavy lines indicate regions of the vectors probed in hybridization analysis. Also shown is the sequences of glZD-GFP (~~SEQ ID NO:1~~ SEQ ID NOS:1-2 and 6, respectively) and glZD-hygro (~~SEQ ID NO:2~~ SEQ ID NOS:7-9, respectively) at locations between *env* gene and IRES, IRES and GFP, and GFP and 3' LTR. Bold letters indicate start or stop codons present within the junctions.